

PF-0621 USN

<110> INCYTE GENOMIC, INC.; TANG, Y. Tom;  
CORLEY, Neil C.; GORGONE, Gina A.;  
GUEGLER, Karl J.; BAUGHN, Mariah R.

<120> MYOSIN HEAVY CHAIN HOMOLOG

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<140> US 09/830,914  
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<150> PCT/US99/26177  
<151> 1999-11-05

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Glu	Arg	Arg	Lys	Tyr	Glu	Gln	Ile	Arg	Asp	Ser	Ile	Ile	Gly	Ile
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Gln	Ala	Met	Phe	Lys	Ala	Asn	Arg	Val	Arg	Arg	Tyr	Val	Glu	Lys
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Ser Pro Glu Phe Leu Val Gln Ile Gln Lys Lys Leu Asn Glu Arg  
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Gly Asn Trp Glu Leu Ala Lys Ile Leu Ser Ile Thr Gly Ser Glu		
80	85	90
Ser Leu Met Ser Leu Ser Glu Glu Lys Val Leu Lys Val Ser Ser		
95	100	105
Asp Ser Leu Leu Pro Ala Asn Pro Glu Ile Leu Asp Gly Val Asp		
110	115	120
Asp Leu Met Gln Leu Ser Tyr Leu Asn Glu Pro Ser Val Leu Tyr		
125	130	135
Asn Leu Gln Tyr Arg Tyr Asp Arg Asp Met Ile Tyr Ser Lys Ala		
140	145	150
Gly Pro Val Leu Val Ala Ile Asn Pro Phe Lys Lys Ile Pro Leu		
155	160	165
Tyr Gly Ser Asp Tyr Ile Glu Ala Tyr Lys Arg Lys Ser Ile Asp		
170	175	180
Asn Pro His Val Tyr Ala Ile Ala Asp Thr Ala Ile Arg Glu Met		
185	190	195
Ile Arg Asp Glu Val Asn Gln Ser Ile Val Ile Ser Gly Glu Ser		
200	205	210
Gly Ala Gly Lys Thr Glu Thr Pro Lys Ile Ala Met Gln Tyr Leu		
215	220	225
Ala Ala Leu Gly Gly Asp Ala Arg Glu Ser Gly Ile Leu Ser		
230	235	240
His Asn Gly Cys Arg Thr Pro Arg Arg Ala Glu Ala Phe Gly Asn		
245	250	255
Ala Lys Thr Ser Arg Asp Asn Asn Ser Ser Arg Ile Gly Lys Leu		
260	265	270
Ile Glu Ile His Phe Ser Glu Thr Gly Lys Ile Ser Gly Ala Lys		
275	280	285
Ile Gln Thr Phe Leu Leu Glu Lys Ser Arg Val Val Gln Cys Thr		
290	295	300
Asp Gly Glu Arg Ser Tyr His Ser Phe Tyr Gln Leu Cys Ala Gly		
305	310	315
Ala Pro Pro Ser Leu Arg Glu Lys Leu Asn Leu Lys Ser Ala Arg		
320	325	330
Glu Tyr Lys Tyr Phe Gln Gln Ser Thr Cys Tyr Ser Ile Asn Gly		
335	340	345
Val Asp Asp Ala Glu Glu Phe Arg Val Val Val Glu Ala Leu Asp		
350	355	360
Ala Val His Val Ser Lys Glu Asn Gln Glu Asn Ala Phe Ala Met		
365	370	375
Leu Ala Ala Val Leu Trp Leu Gly Asn Val Thr Phe Ser Ile Val		
380	385	390
Asp Asn Glu Asn His Val Glu Pro Ile Ile Asp Asp Ala Leu Leu		
395	400	405
Asn Val Ala Lys Leu Ile Gly Cys Glu Ala Asp Asp Leu Lys Leu		
410	415	420
Ala Leu Ser Thr Arg Asn Met Lys Val Gly Asn Asp Ile Ile Val		
425	430	435
Gln Lys Leu Thr Leu Ala Gln Ala Ile Asp Thr Arg Asp Ala Leu		

440	445	450
Ala Lys Ser Ile Tyr Ser Cys Leu Phe Asp Trp Leu Val Glu Gln		
455	460	465
Ile Asn Lys Ser Leu Ala Val Gly Lys Arg Arg Thr Gly Arg Ser		
470	475	480
Ile Ser Ile Leu Asp Ile Tyr Gly Phe Glu Ser Phe Asp Val Asn		
485	490	495
Ser Phe Glu Gln Phe Cys Ile Asn Tyr Ala Asn Glu Arg Leu Gln		
500	505	510
Gln His Phe Asn Arg His Leu Phe Lys Leu Glu Gln Glu Glu Tyr		
515	520	525
Ile Gln Asp Gly Ile Asp Trp Ala Lys Val Asp Phe Glu Asp Asn		
530	535	540
Gln Asp Cys Leu Asn Leu Phe Glu Lys Lys Pro Leu Gly Leu Met		
545	550	555
Thr Leu Leu Asp Glu Glu Ser Thr Phe Pro Asn Gly Thr Asp Met		
560	565	570
Thr Phe Ala Thr Lys Leu Lys Gln His Leu Lys Thr Asn Ser Cys		
575	580	585
Phe Arg Gly Glu Arg Gly Lys Ala Phe Thr Val His His Tyr Ser		
590	595	600
Gly Glu Val Thr Tyr Asp Thr Ser Gly Phe Leu Glu Lys Asn Arg		
605	610	615
Asp Leu Leu His Leu Asp Ser Ile Gln Leu Leu Ser Ser Cys Thr		
620	625	630
Cys Glu Leu Pro Gln Ala Phe Ala Ser Asn Met Leu Ser Leu Ser		
635	640	645
Glu Lys Pro Val Pro Gly Pro Leu His Lys Ser Gly Gly Ala Asp		
650	655	660
Ser Gln Lys Leu Ser Val Val Thr Lys Phe Lys Gly Gln Leu Phe		
665	670	675
Gln Leu Met Gln Arg Leu Glu Ser Thr Thr Pro His Phe Ile Arg		
680	685	690
Cys Ile Lys Pro Asn Asn Ser Gln Ser Pro Gly Ile Tyr His Gln		
695	700	705
Gly Leu Val Leu Gln Gln Leu Arg Cys Cys Gly Val Leu Glu Val		
710	715	720
Val Arg Ile Ser Arg Ser Gly Phe Pro Thr Arg Met Ser His Gln		
725	730	735
Lys Phe Ala Arg Arg Tyr Gly Phe Leu Leu Leu Glu His Val Ala		
740	745	750
Ser Gln Asp Pro Leu Ser Val Ser Val Ala Ile Leu His Gln Phe		
755	760	765
Asp Ile Leu Pro Glu Met Tyr Gln Ile Gly Tyr Thr Lys Leu Phe		
770	775	780
Phe Arg Thr Gly Gln Ile Gly Lys Leu Glu Asp Thr Arg Asn Arg		
785	790	795
Thr Leu Asn Gly Ile Leu Arg Val Gln Ser Cys Phe Arg Gly His		
800	805	810
Lys Ala Arg Gln Tyr Met Lys Glu Leu Lys Arg Gly Ile Phe Asn		
815	820	825
Leu Gln Ala Phe Ala Arg Gly Glu Lys Thr Arg Lys Glu Phe Ala		
830	835	840
Ile Leu Val His Arg His Arg Ala Ala Val His Ile Gln Lys His		

845	850	855
Ile Lys Ala Lys Ile Ser Lys Lys Arg Phe Glu Asp Val His Gly		
860	865	870
Ala Thr Ile Thr Leu Gln Ala Val Ile Arg Gly Trp Leu Val Arg		
875	880	885
Arg Cys Ser Gly Asp Ile Ala Leu Leu Gln Phe Gly Ser Gly Lys		
890	895	900
Gly Asn Gly Ser Asp Glu Val Leu Val Lys Ser Ser Tyr Leu Ala		
905	910	915
Glu Leu Gln Arg Arg Ile Leu Lys Ala Glu Ala Gly Leu Arg Glu		
920	925	930
Lys Glu Glu Glu Asn Asp Ile Leu His Gln Arg Leu Gln Gln Tyr		
935	940	945
Glu Asn Arg Trp Ser Glu Tyr Glu Leu Lys Met Lys Ser Met Glu		
950	955	960
Glu Val Trp Gln Lys Gln Met Arg Ser Leu Gln Ser Ser Leu Ser		
965	970	975
Ile Ala Lys Lys Ser Leu Ser Tyr Asp Asp Ser Glu Arg Asn Ser		
980	985	990
Asp Ala Ser Ile Asn Thr Ala Asn Asp Glu Thr Asn Pro Pro Trp		
995	1000	1005
Asp Ala Ala Thr Asn Gly Arg Arg Asn Gly Val Glu Asn Val Arg		
1010	1015	1020
Pro Met Ser Ala Gly Leu Ser Val Ile Ser Arg Leu Ala Glu Glu		
1025	1030	1035
Phe Glu Gln Arg Ser Gln Val Phe Gly Asp Asp Ala Lys Phe Leu		
1040	1045	1050
Val Glu Val Lys Ser Gly Gln Val Glu Ala Asn Leu Asn Pro Asp		
1055	1060	1065
His Glu Leu Arg Arg Leu Lys Gln Met Phe Glu Gly Trp Lys Lys		
1070	1075	1080
Asp Tyr Thr Ala Arg Leu Arg Glu Thr Lys Val Ile Leu Asn Lys		
1085	1090	1095
Leu Gly His Glu Asp Gly Asp Gly Glu Lys Gly Lys Lys Lys Trp		
1100	1105	1110
Trp Gly Arg Leu Asn Ser Ser Arg Val Asn		
1115	1120	